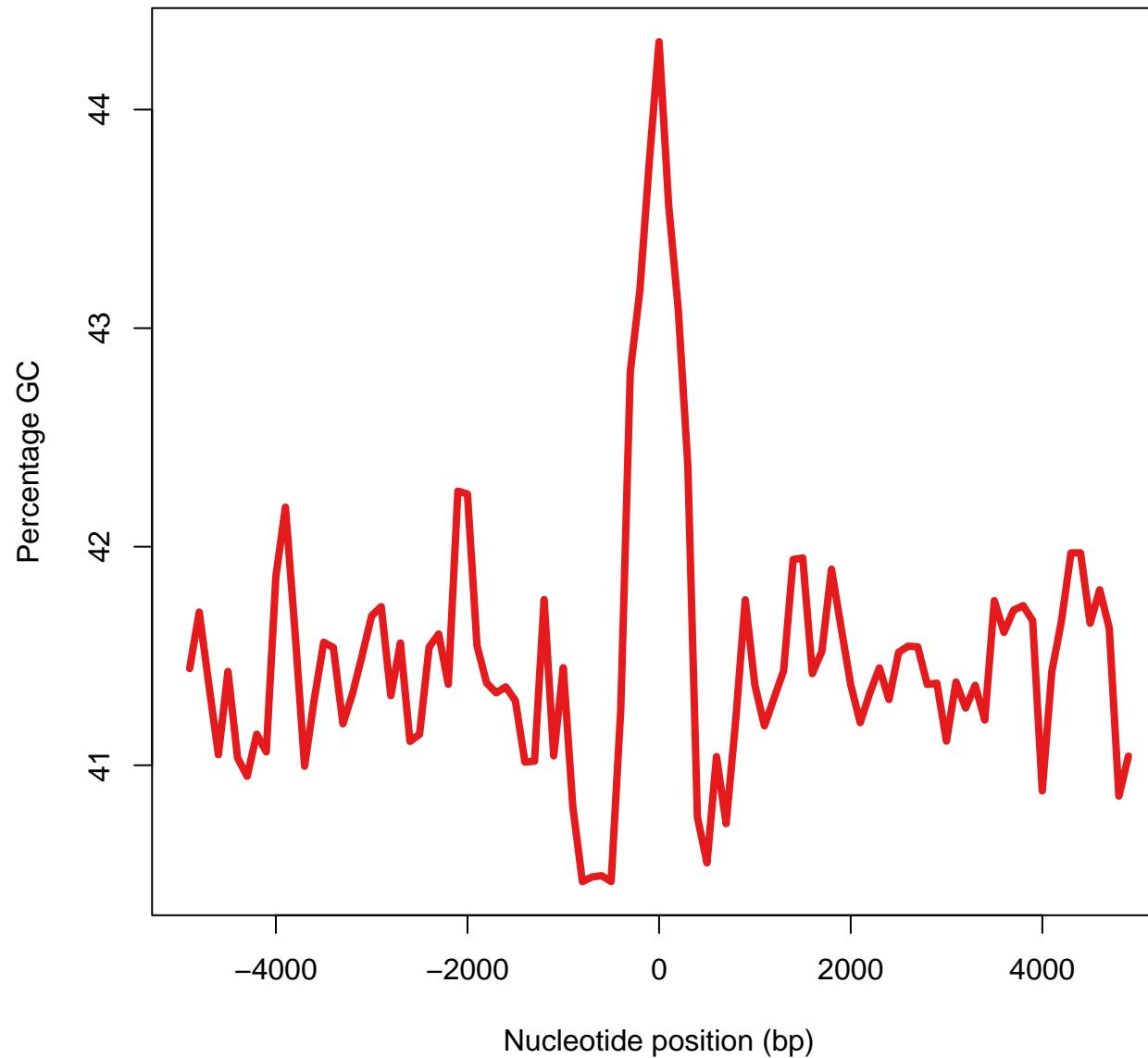
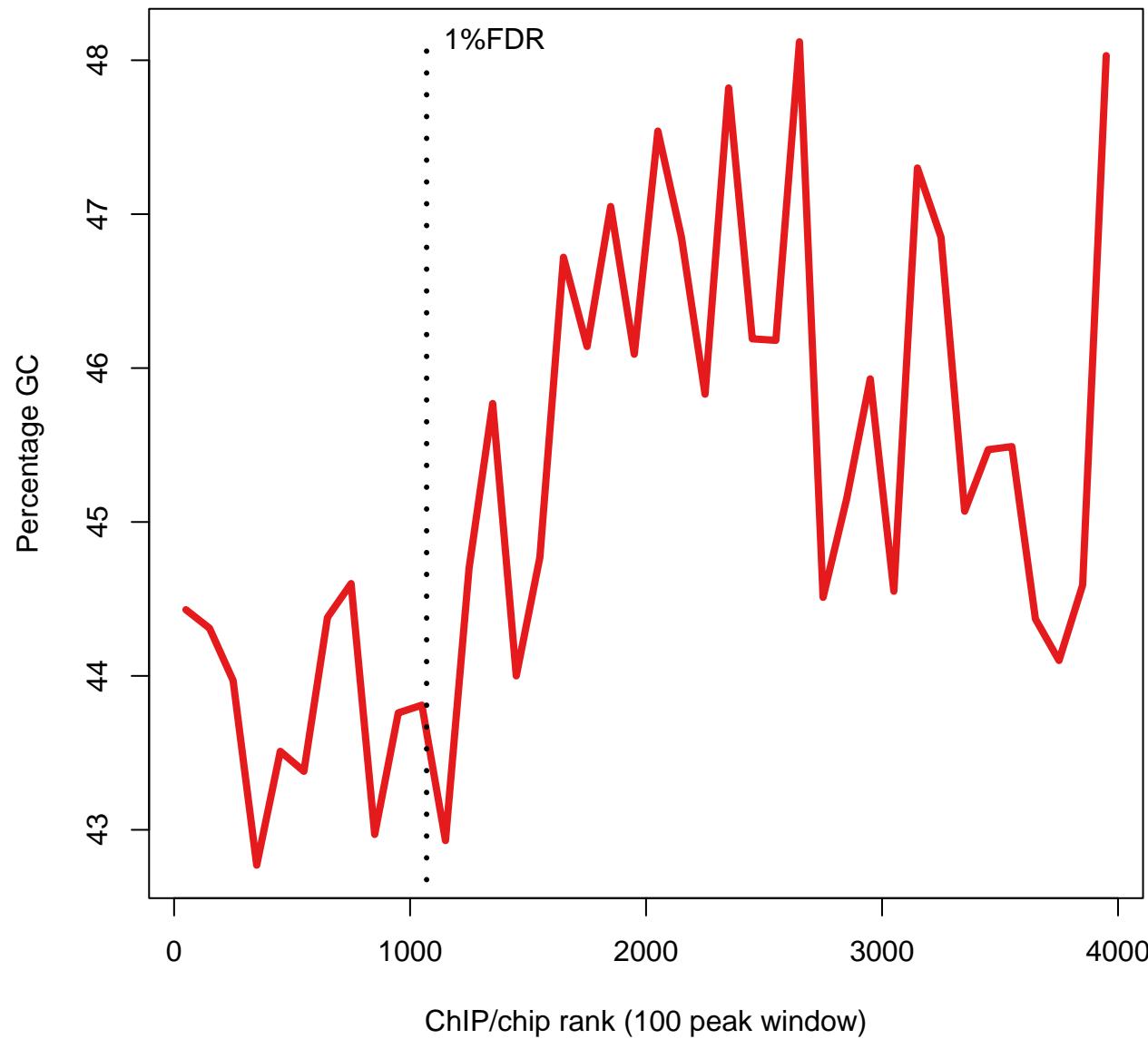


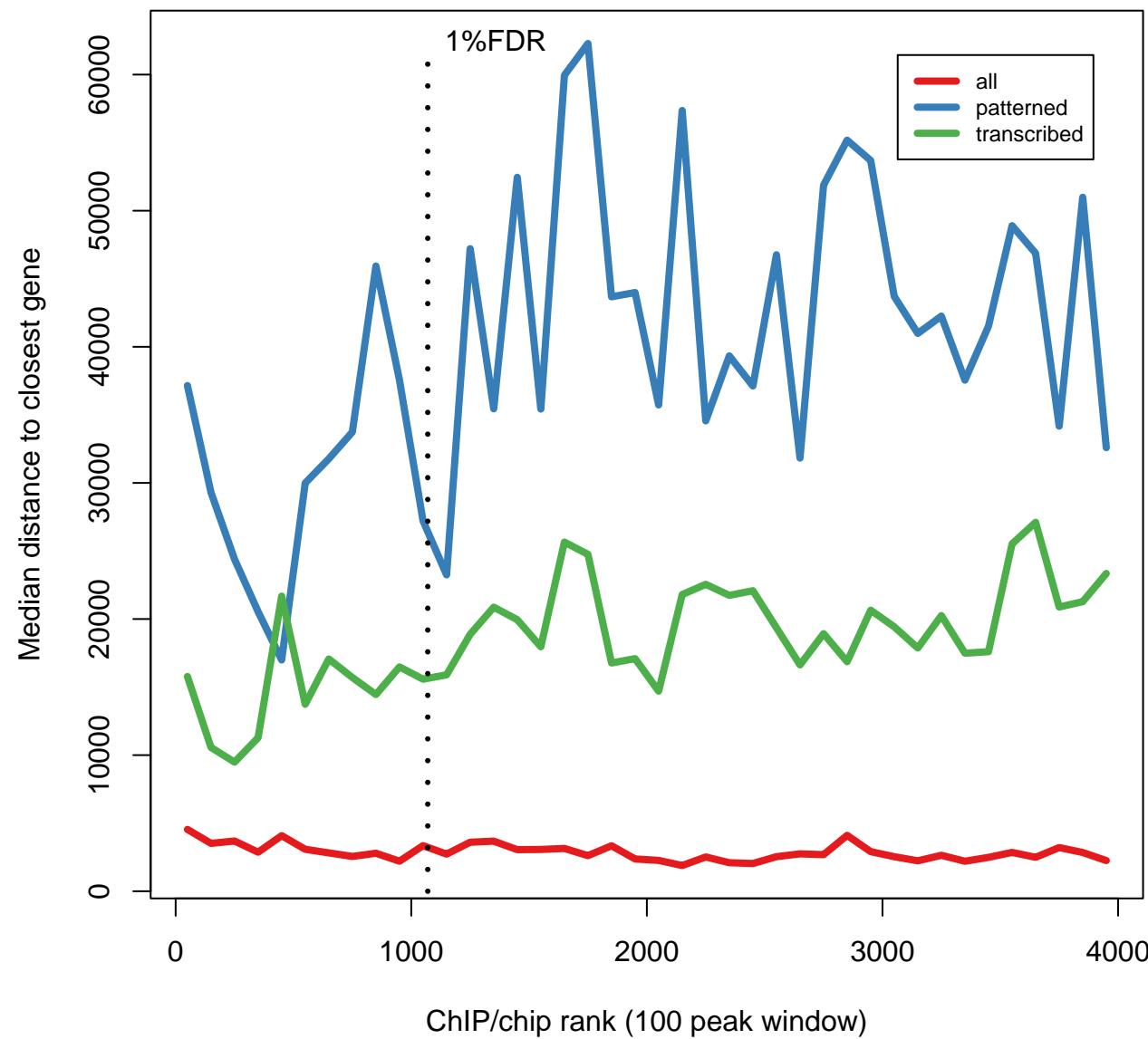
gt_2 base composition across peaks



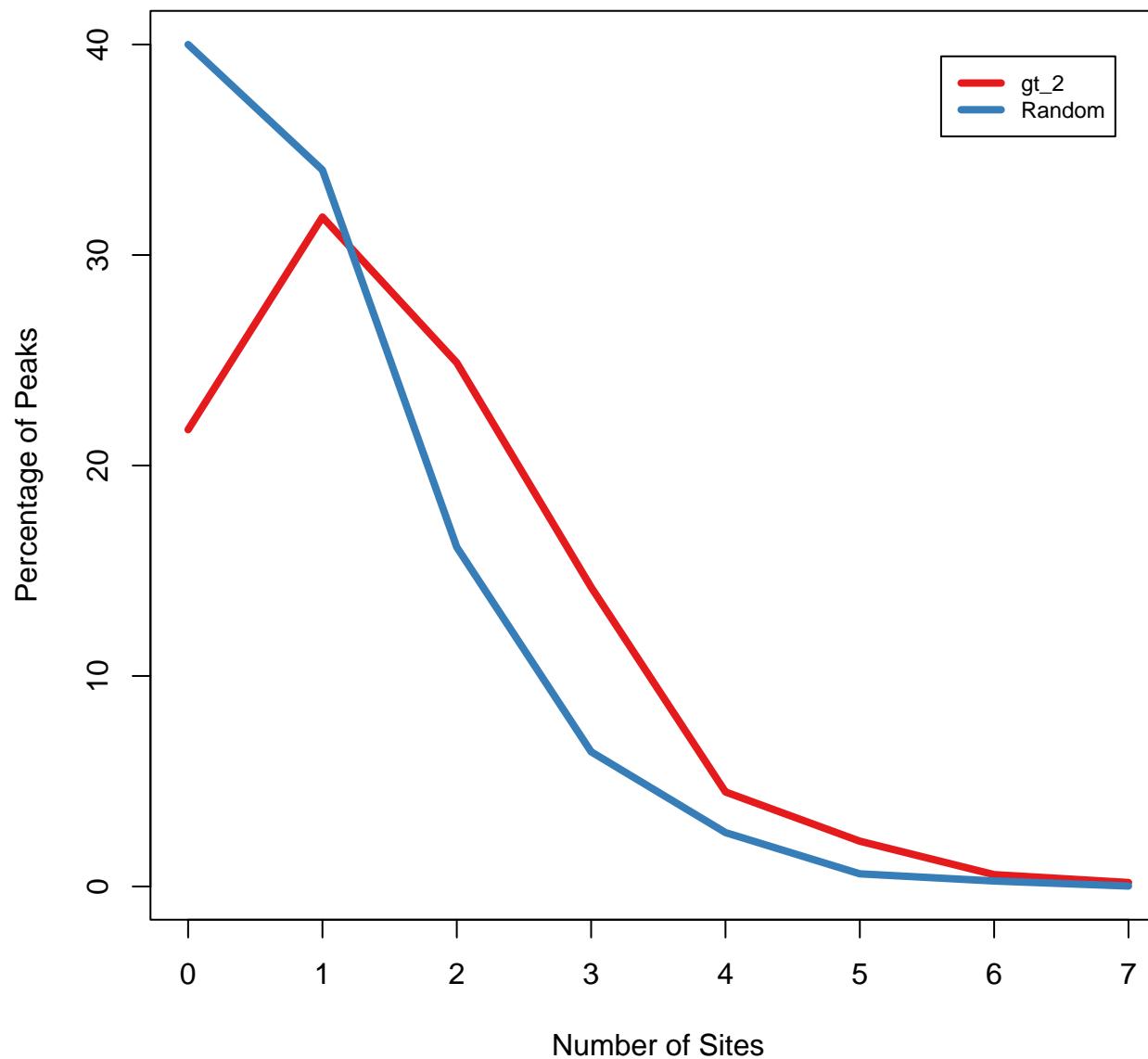
gt_2 base composition down ranks



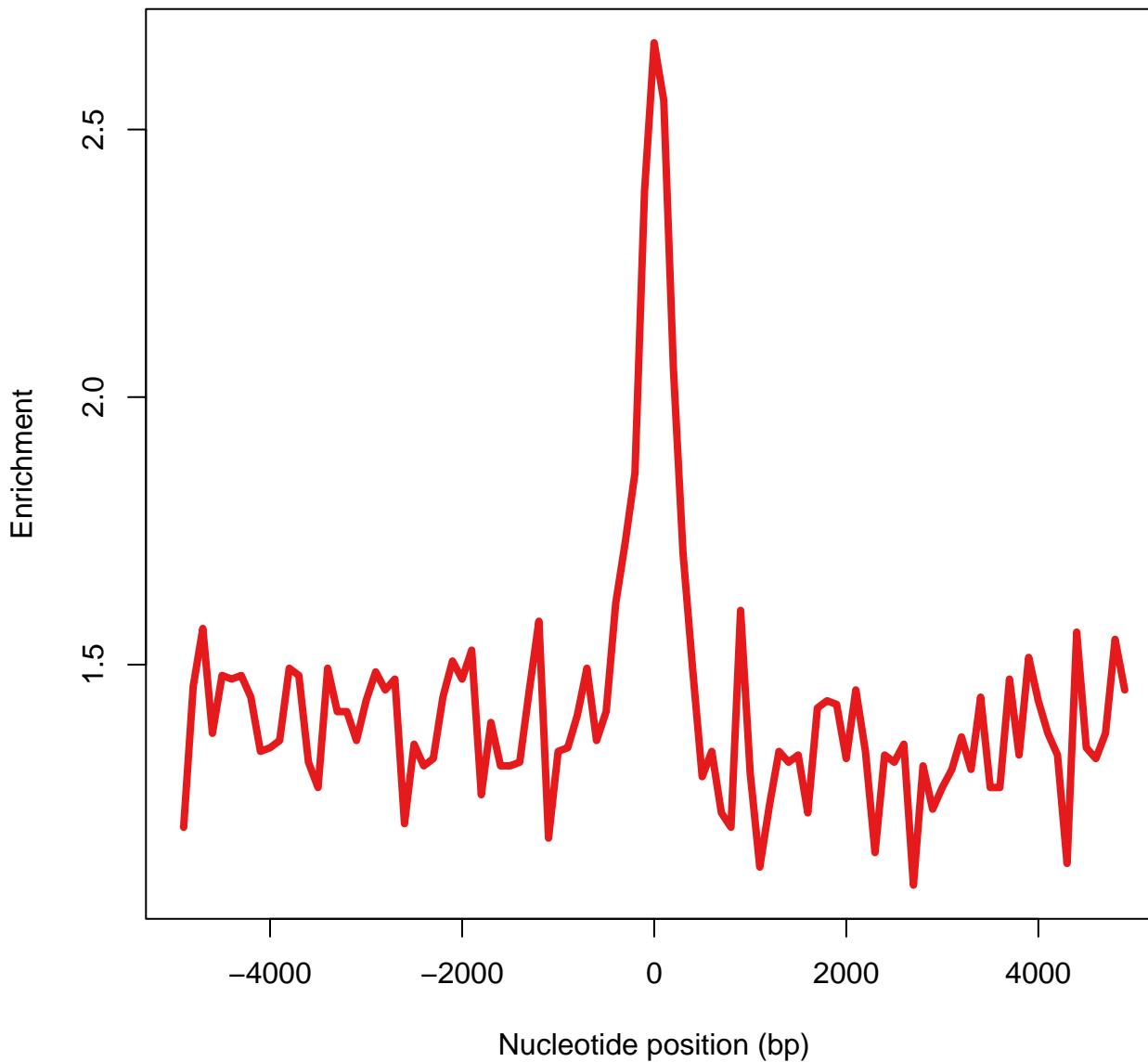
gt_2 median distance to genes



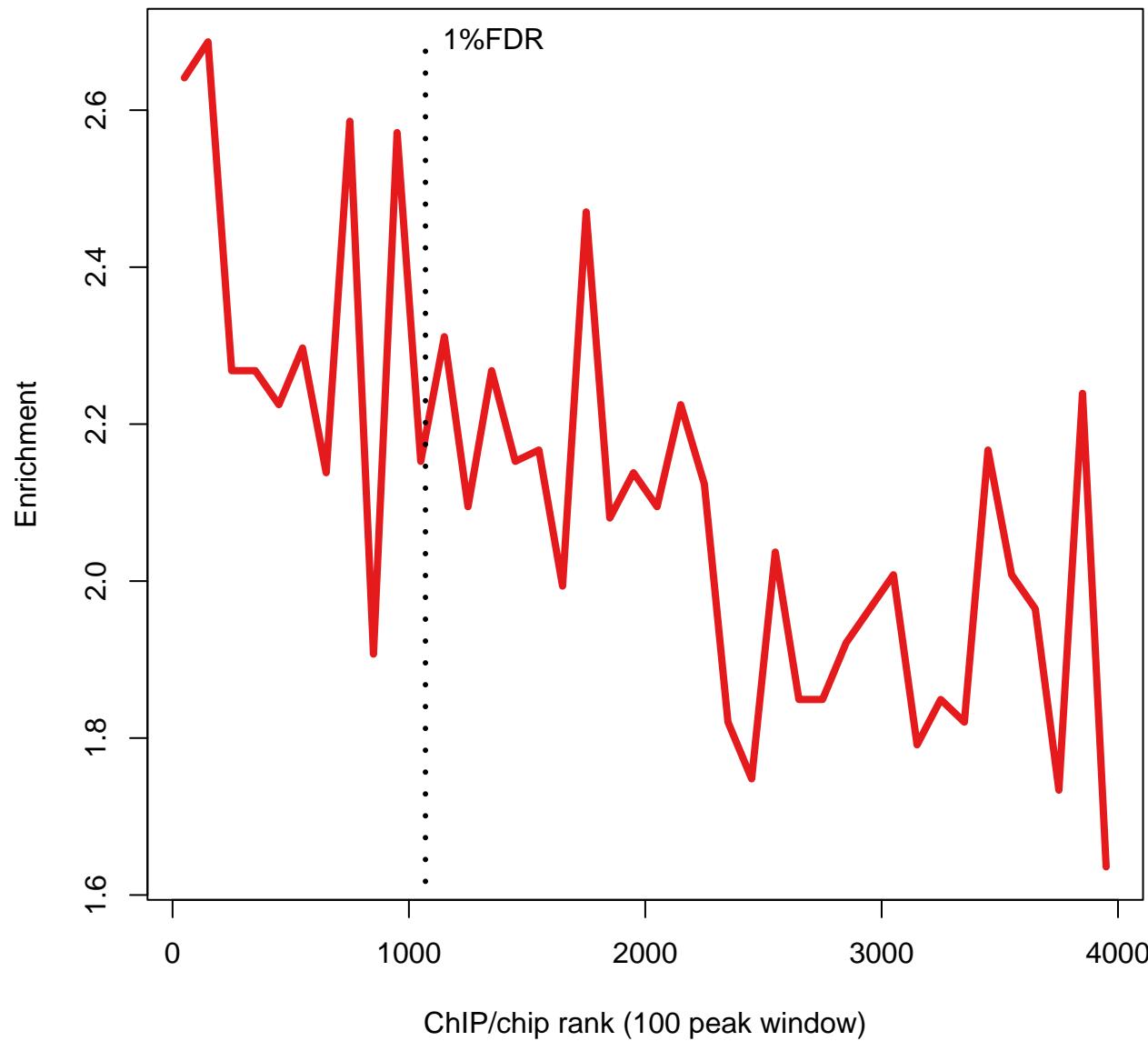
gt_2 distribution of number of sites



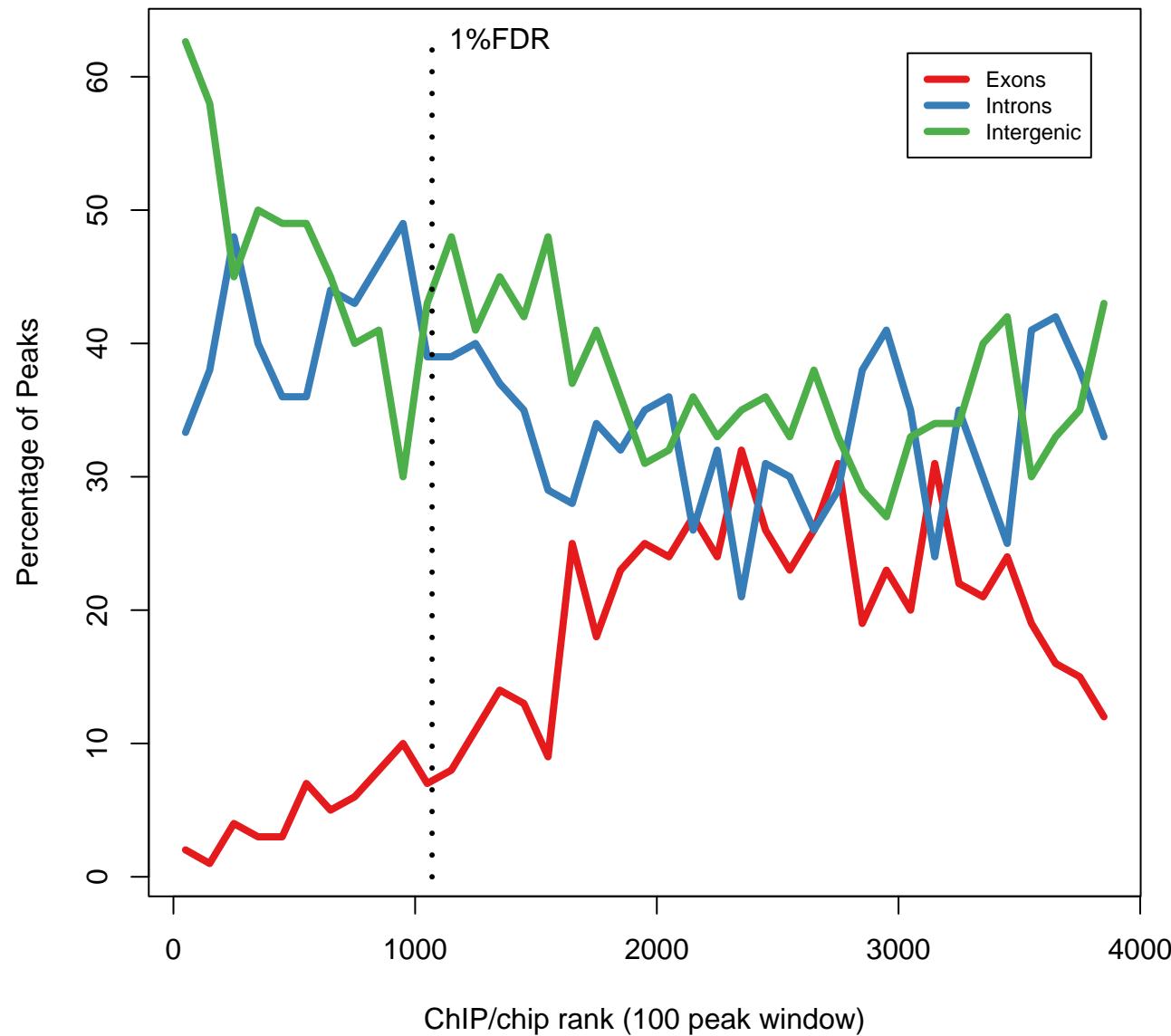
gt_2 PWM enrichment across peaks



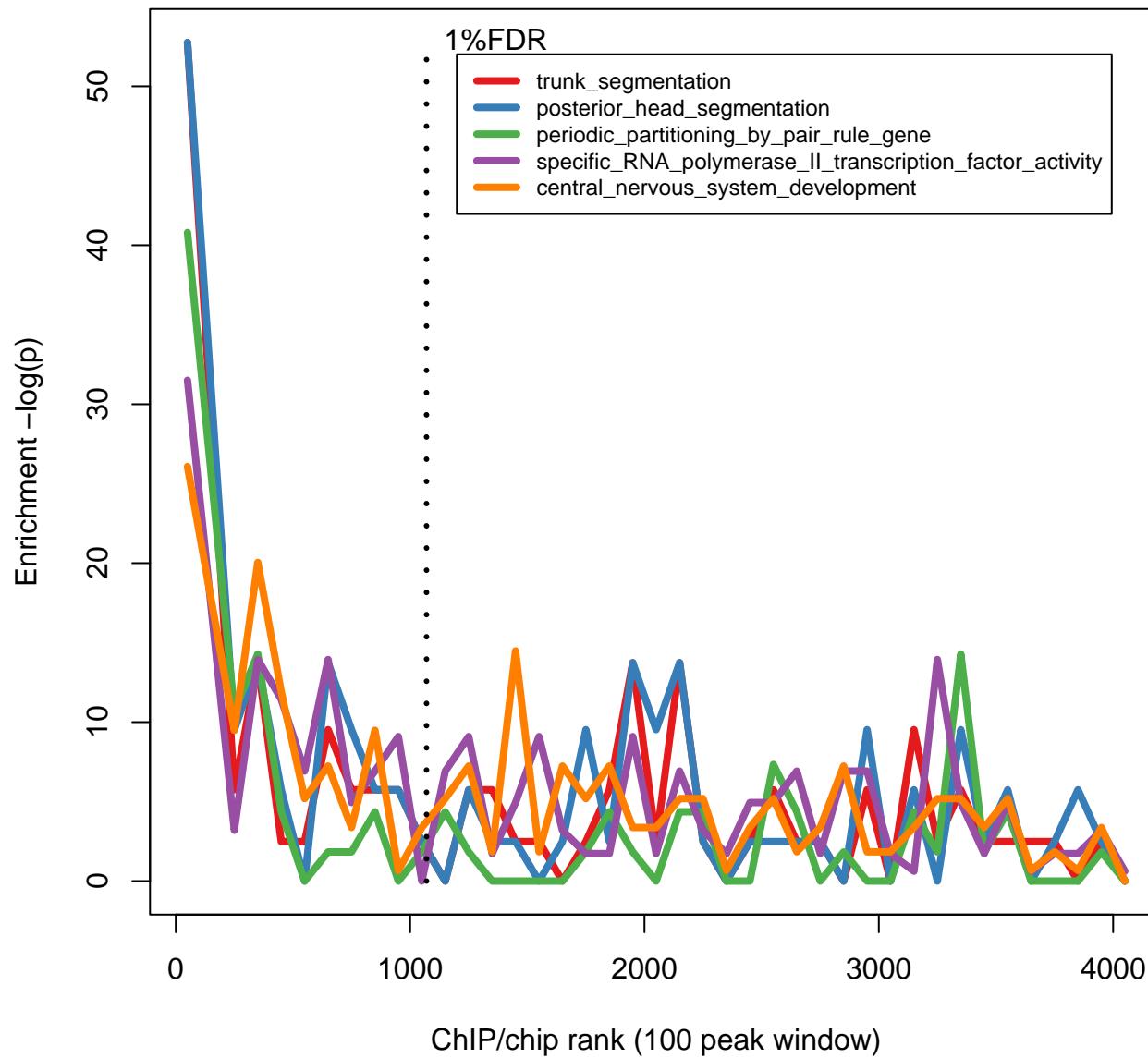
gt_2 PWM enrichment down ranks

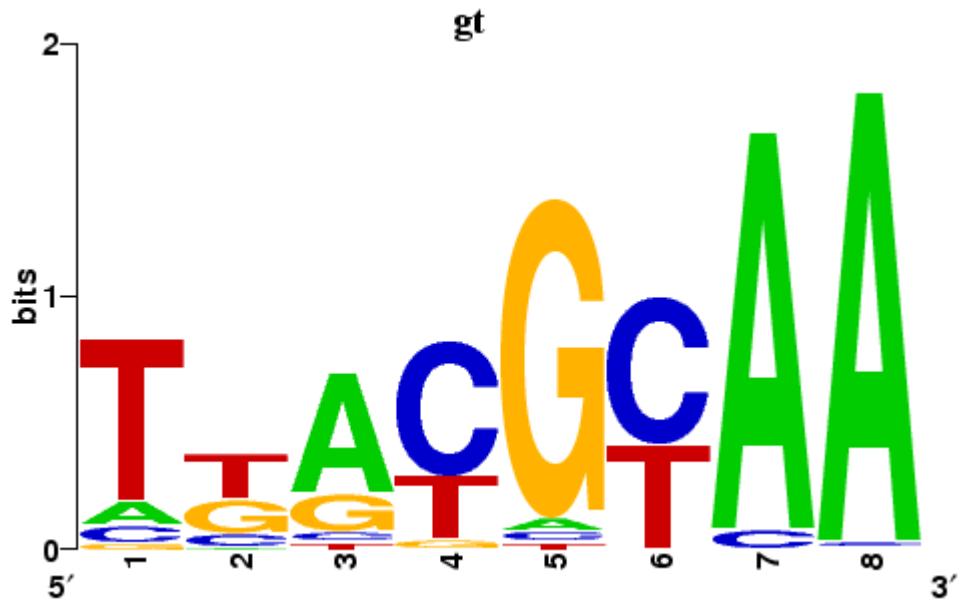


gt_2 genomic location

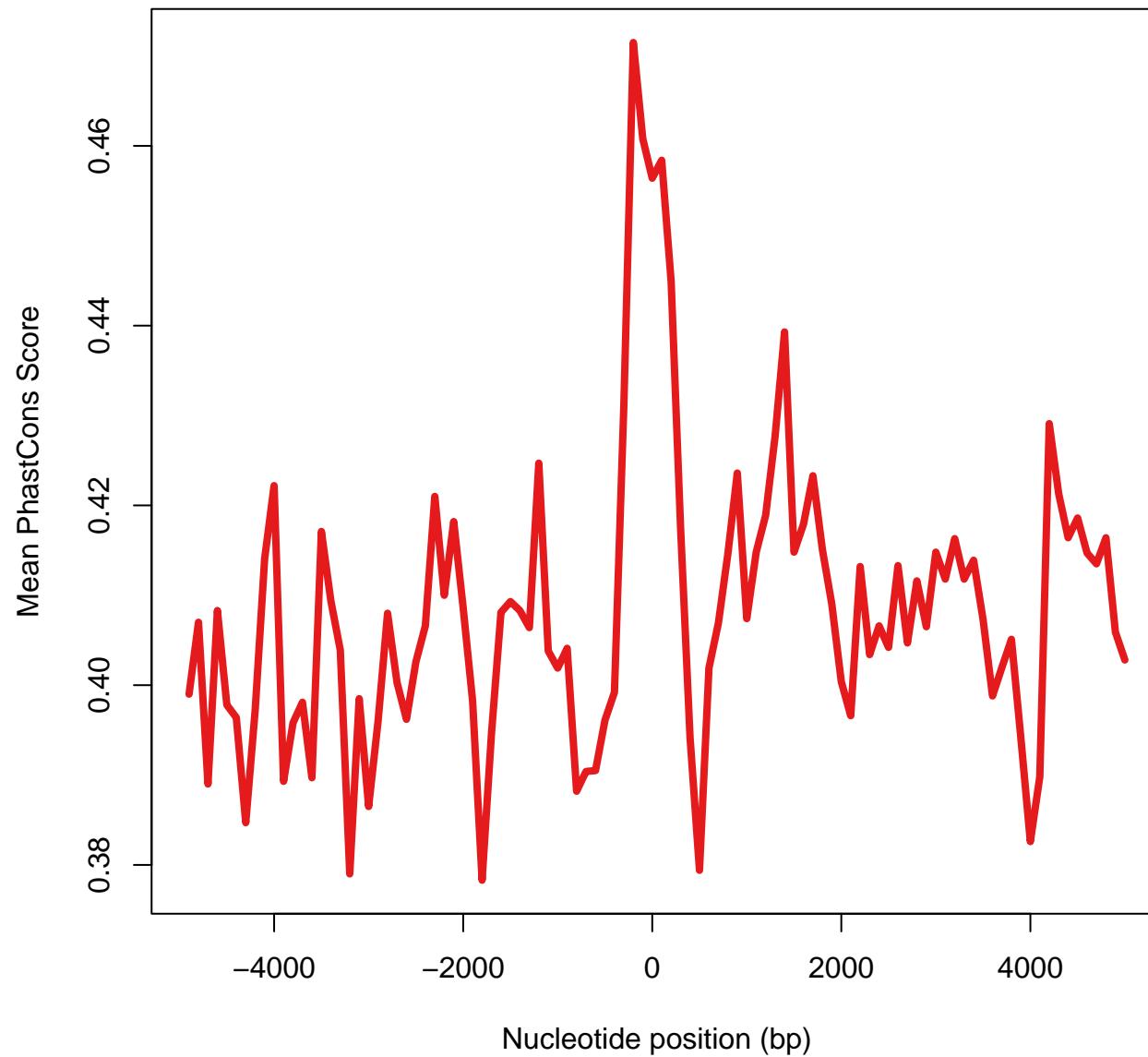


gt_2 GO term enrichment (-logP)





gt_2 mean PhastCons across peaks



gt_2 mean PhastCons down ranks

